

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
1 May 2003 (01.05.2003)

PCT

(10) International Publication Number
WO 03/035110 A1

(51) International Patent Classification⁷: **A61K 45/00**,
A61L 31/16, A61P 9/10, A61K 31/7105, 38/17, 39/395

(21) International Application Number: PCT/US02/34120

(22) International Filing Date: 23 October 2002 (23.10.2002)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
60/335,637 24 October 2001 (24.10.2001) US
60/341,359 17 December 2001 (17.12.2001) US
10/128,166 23 April 2002 (23.04.2002) US

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZM, ZW.

(84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

(71) Applicant: **CEDARS-SINAI MEDICAL CENTER**
[US/US]; 8700 Beverly Boulevard, Los Angeles, CA 90048 (US).

(72) Inventors: **ARDITI, Moshe**; 4210 Hayvenhurst Avenue, Encino, CA 91436 (US). **RAJAVASHISTH, Tripathi**; 15203 Florwood Avenue, El Camino Village, CA 90260 (US). **SHAH, Prediman, K.**; 111 North Layton Drive, Los Angeles, CA 90049 (US).

(74) Agents: **LEVY, Seth, D.** et al.; Pillsbury Winthrop Llp, 725 South Figueroa Street, Suite 2800, Los Angeles, CA 90017-5406 (US).

Published:

- with international search report
- before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: TREATING VASCULAR DISEASE BY INHIBITING TOLL-LIKE RECEPTOR-4

(57) Abstract: Methods included herein describe the treatment of atherosclerosis and other vascular diseases such as thrombosis, restenosis after angioplasty and/or stenting, and vein-graft disease after bypass surgery, by inhibition of the expression or biologic activity of Toll-like receptor-4 (TLR-4). Also included is an intravascular device coated with a compound that inhibits TLR-4; thereby imparting an improved efficacy to the device. TLR-4 cell signal transduction is at least partially responsible for the manifestation, continuation, and/or worsening of atherosclerosis and other forms of vascular disease. The present invention provides several means with which to inhibit this signal transduction pathway.

WO 03/035110 A1

TREATING VASCULAR DISEASE BY INHIBITING TOLL-LIKE RECEPTOR-4

FIELD OF THE INVENTION

This invention relates to methods for inhibiting the biological activity of Toll-like
5 receptor-4 ("TLR-4"), and, in particular, to methods for treating vascular disease by inhibiting the
expression or signaling by TLR-4.

BACKGROUND OF THE INVENTION

Heart disease remains the leading cause of death worldwide, accounting for nearly 30% of
10 the annual total (i.e., approximately 15 million people). Heart and vascular disease debilitate
many more individuals every year. For many, atherosclerotic disease is a life-long process; it may
possess an initial stage in childhood, without clinical manifestation until middle age or later. Its
development has been repeatedly linked to unhealthy lifestyles (e.g., tobacco use, unbalanced diet,
and physical inactivity). Much progress has been made in the detection and treatment of various
15 forms of heart and vascular disease, but preventative measures and assorted treatment regimens
are usually incapable of halting or curing the underlying disease condition.

Experimental work over the past decade has linked inflammation of the blood vessel wall
to atherogenesis, restenosis, and plaque disruption. The precise triggers for inflammation are not
known, but it is believed that some triggers may include modified lipoproteins and various local
20 or distant infections. A potential role for infection in the development of atherosclerosis has been
considered; specific infectious agents, such as *Chlamydia pneumoniae* ("C. pneumoniae"), have
been suggested as playing a role in the progression and/or destabilization of atherosclerosis.

Recent studies suggest that chlamydia lipopolysaccharide ("cLPS") induces foam-cell
formation, whereas its heat-shock protein ("cHSP-60") induces oxidative modification of low-
25 density lipoproteins ("LDL"). M.V. Kalayoglu and G.I. Byrne, "*Chlamydia pneumoniae*

component that induces macrophage foam cell formation is chlamydial lipopolysaccharide,"

Infect. & Immunity 66:5067-5072 (1998); G.I. Byrne and M.V. Kalayoglu, "*Chlamydia*

pneumoniae and atherosclerosis: Links to the disease process," *Amer. Heart Journal* 138:S488-

S490 (1999). cHSP-60 has been implicated in the induction of deleterious immune responses in

5 human chlamydial infection and has been found to co-localize with infiltrating macrophages in

atheroma lesions. A.G. Kol *et al.*, "Chlamydial heat shock protein 60 localizes in human

atheroma and regulates macrophage tumor necrosis factor alpha and matrix metalloproteinase

expression," *Circulation* 98:300 (1998). Collectively, these data support a potential role for *C.*

pneumoniae in the development and progression of atherosclerosis and suggest that this organism

10 may indeed play an active role in atheroma development. However, available data underscore the

current lack of an understanding of the molecular mechanisms that link *C. pneumoniae* infections

to innate immunity and trigger the signals for enhanced inflammation and atherogenesis. Absent

such an understanding, it is quite difficult to develop a useful mechanism for treating vascular

disease based on these data.

15 Although precise triggers for inflammation in atherosclerosis are not fully understood,

hypercholesterolemia, modified lipoproteins, and infection with organisms such as *C. pneumoniae*

and others have been implicated. There is evidence that *C. pneumoniae* infection can accelerate

the progression and facilitate the induction of atherosclerosis in cholesterol-fed rabbits and

genetically modified atherosclerosis prone mice. Without a clear understanding of the mechanism

20 that controls this system, however, these data may not provide the basis for a treatment or cure for

atherosclerosis. J.B. Muhlestein *et al.*, "Infection with *Chlamydia pneumoniae* accelerates the

development of atherosclerosis and treatment with azithromycin prevents it in a rabbit model,"

Circulation 97:633-636 (1998); T.C. Moazed *et al.*, "Murine models of *Chlamydia pneumoniae*

infection and atherosclerosis," *J. Infect. Dis.* 175:883-890 (1997); T.C. Moazed *et al.*, "*Chlamydia*

25 *pneumoniae* infection accelerates the progression of atherosclerosis in Apolipoprotein E-deficient

mice," *J. Infect. Dis.* 180:238-241 (1999); L.A. Campbell and C.C. Kuo, "Mouse models of *Chlamydia pneumoniae* infection and atherosclerosis," *Am. Heart J.* 138:S516-S518 (1999); K. Laitinen *et al.*, "*Chlamydia pneumoniae* infection induces inflammatory changes in the aortas of rabbits," *Infect & Immunity* 65:4832-4835 (1997).

5 The concept of *C. pneumoniae*-induced atherogenesis is strengthened by the finding that antibiotic therapy against chlamydia prevents acceleration of atherosclerosis in the rabbit model. Ingalls *et al.* have suggested lipopolysaccharide ("LPS"), and Kol *et al.* have implicated HSP-60 as the triggers for chlamydia-induced inflammatory responses. R.R. Ingalls *et al.*, "The inflammatory cytokine response to *Chlamydia trachomatis* infection is endotoxin mediated,"
10 *Infect & Immun.* 63:3125-3130 (1995); A. Kol *et al.*, "Chlamydial and human heat shock protein 60s activate human vascular endothelium, smooth muscle cells and macrophages," *J Clin Invest* 103:571-577 (1999); A. Kol *et al.*, "Heat shock protein (HSP)60 activates the innate immune response," *The J of Immunol.* 164:13-17 (2000). To date, however, the precise molecular mechanisms by which infections such as *C. pneumoniae* contribute to the progression of
15 atherosclerosis and the links among lipids, microbial antigens, and innate immune and inflammatory responses are not well understood.

One recent study, however, indicated that HSP-60 induces smooth muscle cell proliferation *in vitro*; smooth muscle cell proliferation being directly related to atherogenesis. Sasu *et al.*, "Chlamydia pneumoniae and Chlamydial Heat Shock Protein 60 Stimulate
20 Proliferation of Vascular Smooth Muscle Cells via Toll-Like Receptor 4 and p44/p42 Mitogen-Activated Protein Kinase Activation," *Circ. Res.* 89:244-250 (2001). The study showed that smooth muscle cell proliferation was blocked or severely hampered by anti-TLR-4 antibodies. This finding suggests that HSP-60 also causes smooth muscle cell proliferation via a TLR-4 pathway.

The introduction of surgical and percutaneous arterial revascularization to treat atherosclerosis has profoundly altered the clinical management of disease, but has also brought unanticipated problems and unanswered questions. Surgical, and especially percutaneous revascularization, may elicit an exaggerated healing response, which in many respects is similar to the development of *de novo* atherosclerotic lesions. This "response to injury" is more proliferative in nature than *de novo* lesion formation, but may nevertheless lead to restenosis, or even late or abrupt vessel closure, and may ultimately result in a failed revascularization attempt. For this and additional reasons, long-term clinical studies have documented improved outcomes only in select patient subgroups; for those with stable angina pectoris, coronary intervention remains merely palliative, and does not alter the progression or outcome of the underlying causative disease process.

With balloon coronary angioplasty, restenosis rates of 30%-40% or more have been documented, and certain lesion sites and patient subgroups have been found to be particularly susceptible to restenosis. Intensive research efforts into the cause of restenosis have yielded considerable insight, but as yet no unequivocal treatment has been identified to eliminate the problem. Technical innovations in revascularization equipment and techniques have shown some success, but even this has been of limited efficacy. In particular, the development of the intracoronary stent markedly reduced the incidence of restenosis. With proper stent placement techniques, restenosis rates have been reduced to roughly 15%-30%, so intracoronary stent placement has largely supplanted balloon angioplasty alone as the interventional coronary treatment of choice. Still, given the rapid proliferation and acceptance of intracoronary stenting, even a 15%-30% restenosis rate results in a very large number of patients in whom the revascularization attempt has been unsuccessful, and for whom other treatment strategies have not been sufficiently effective. Often, the same patient may need multiple separate interventions, and ultimately these may not be successful.

Since the arterial response to injury is predominantly mitogenic and neoproliferative in nature, intracoronary irradiation (or intracoronary brachytherapy) has been developed and deployed to attempt to reduce further the number of patients who restenose following coronary intervention. Intracoronary brachytherapy has also met with limited success, however, and has brought with it two new manifestations of the disease as a side effect: geometric miss and late in-stent thrombosis. It appears likely that these two effects will significantly limit the efficacy of intracoronary brachytherapy as a definitive treatment for restenosis. Thus, a need remains for an effective way to limit or eliminate restenosis following coronary stent placement. Alternatively, if intracoronary brachytherapy is to achieve unequivocal effectiveness in eliminating restenosis following stent placement, a solution to late in-stent thrombosis and geometric miss must be found.

Conventional treatments for vascular disease have substantial drawbacks; many are only partially effective, and few provide a true cure for associated conditions. There remains a clear need in the art for a method of preventing, treating, and curing vascular disease, including atherosclerosis. There remains a further need in the art for improvements to present stent technology, whereby one can minimize the chance of restenosis.

SUMMARY OF THE INVENTION

It is an object of the present invention to provide methods for inhibiting the biological activity of TLR-4, as, for example, by inhibiting its expression or signaling. It is a further object of the invention to provide methods of treating those diseases in which inhibiting TLR-4 would have a beneficial effect. Such diseases include, for example, vascular disease such as atherosclerosis and thrombosis, restenosis after angioplasty and/or stenting, and vein-graft disease after bypass surgery.

A first embodiment of the invention is directed to a method of inhibiting TLR-4 by administering to a mammal recombinant viral vectors (e.g., adenovirus, adeno-associated virus, retroviruses, lentiviruses, or other viral vectors) that deliver genes expressing antisense TLR-4 RNA; doing so inhibits the expression of TLR-4, thereby inhibiting its biological activity. An
5 optimal amount of viral particles and an effective and convenient route to administer it (e.g., by administering it intravenously or intramuscularly) can readily be determined by one of ordinary skill in the art of microbiology.

A second embodiment of the present invention is directed to a method of inhibiting TLR-4 signaling by inducing *in vivo* production of a high affinity soluble TLR-4 protein that competes
10 for non-bound HSP-60, LPS, and other ligands that are molecularly configured to operatively interact with a TLR-4 receptor. The TLR-4 protein most preferably lacks the TLR-4 signal transduction domain, or at least a sufficient amount of the TLR-4 signal transduction domain such that the TLR-4 protein is unable to participate in TLR-4 signal transduction. The method involves delivering viral vectors to produce an amount of soluble TLR-4 or its derivatives that is sufficient
15 to reduce the amount of HSP-60, LPS, or other ligands that are molecularly configured to operatively interact with a TLR-4 receptor, thereby inhibiting the TLR-4 signaling pathway.

A third embodiment of the present invention is directed to a method of inhibiting TLR-4 signaling with somatic-cell gene therapy. According to this method, one administers a ribozyme-viral (adeno, adeno-associated, lentiviral or other) vector against TLR-4 mRNA in a mammal.
20 The method utilizes a hammerhead ribozyme expression cassette in a viral backbone. Ribozymes have sequence-specific endoribonuclease activity, which makes them useful for sequence-specific cleavage of mRNAs and further inhibition of gene expression. Ribozyme therapy is widely regarded as a new and potential pharmaceutical class of reagent to treat a number of medical disorders. A desired quantity or the length of expression of the ribozyme-viral vector can be
25 readily determined without undue experimentation, as can the most effective and convenient route

of administering it. Ribozyme-viral vectors against TLR-4 mRNA permit one to uniquely assess the contribution of TLR-4 mediated cell-signaling to vascular physiology, and to therapeutically intervene in the pathology such signaling causes.

A fourth embodiment of the present invention provides a non-viral method to inhibit the
5 expression of TLR-4. This method involves antisense therapy using oligodeoxynucleotides (“ODN”) that inhibit the expression of the TLR-4 gene product by specific base pairing of single stranded regions of the TLR-4 mRNA. The method involves synthesis of ODN complimentary to a sufficient portion of TLR-4 mRNA. The method further provides an effective amount of ODN to inhibit the TLR-4 signaling pathways in a mammal.

10 A fifth embodiment of the present invention provides a method to inhibit the expression of TLR-4 by RNA interference (“RNAi”). This method involves the use of double-stranded RNA (“dsRNA”) that are sufficiently homologous to a portion of the TLR-4 gene product such that the dsRNA degrades mRNA that would otherwise affect the production of TLR-4. A well-defined 21-base duplex RNA, referred to as small interfering RNA (“siRNA”), may operate in
15 conjunction with various cellular components to silence the TLR-4 gene product with sequence homology.

A sixth embodiment of the present invention provides a method to inhibit the TLR-4 cell-signaling pathway by peptide mimetics. This method involves the introduction of small peptides (i.e., peptides of approximately 10-20 amino acids) that bind to TLR-4 ligands, thereby preventing
20 proper TLR-4 ligands from binding to TLR-4 or associated receptors (e.g., MD2). In this manner, the TLR-4 cell-signaling pathway may be blocked from signal transduction, because the proper TLR-4 ligands are unable to bind correctly to TLR-4 or its associated receptors.

A seventh embodiment of the present invention provides a method to inhibit the expression of TLR-4 through the introduction an anti-TLR-4 antibody. Such an antibody may be
25 delivered to a mammal through any conventional mechanism in an amount effective to inhibit the

TLR-4 signaling pathways in a mammal; the mechanism of delivery and quantity of antibody necessary for inhibiting TLR-4 expression both being readily ascertainable without undue experimentation.

Other features and advantages of the invention will become apparent from the following detailed description, taken in conjunction with the accompanying drawings, which illustrate, by way of example, various features of embodiments of the invention.

BRIEF DESCRIPTION OF THE FIGURES

The file of this patent contains at least one drawing executed in color. Copies of this patent with color drawing(s) will be provided by the Patent and Trademark Office upon request and payment of the necessary fee.

Fig. 1 is executed in color. Fig. 1a is a histologic depiction of TLR-4 immunoreactivity (brown) within the lipid core of an atherosclerotic plaque in the aortic sinus of an apolipoprotein E-deficient mouse. Figs. 1b and 1c depict the histology of macrophage (brown) and smooth muscle cell (red) immunoreactivity, respectively, in the serial section of the same aortic sinus. Fig. 1d depicts Rabbit IgG staining for a negative control. Fig. 1e depicts a lack of TLR-4 immunoreactivity in the non-atherosclerotic aortic mouse sinus.

Fig. 2 is executed in color, and is a series of photomicrographs indicating TLR-4 expression in human atherosclerotic lipid-rich plaques, and a lack of such expression in fibrous plaques. Fig. 2a depicts an atherosclerotic plaque stained brown with rabbit anti-human TLR-4 antiserum. Fig 2b depicts a negative control where the primary antibody was replaced by rabbit IgG. Fig 2c depicts TLR-4 immunoreactivity (brown). Fig. 2d depicts a double immunostain of TLR-4 (brown) and macrophages (red), demonstrating co-localization. Fig. 2e depicts macrophage immunoreactivity (red), under a higher magnification. Fig. 2f depicts TLR-4 immunoreactivity (brown), under a higher magnification. Fig. 2g depicts macrophage (red) along

with TLR-4 (brown) immunoreactivity, under a higher magnification. Fig. 2h depicts a lack of immunoreactivity of TLR-4 in a fibrous plaque. Fig. 2i depicts smooth muscle cell alpha actin immunoreactivity (red) without TLR-4 immunoreactivity (brown) upon double-staining. Fig. 2j depicts a lack of immunoreactivity of macrophages in a fibrous plaque. Fig. 2k depicts a negative control using pre-absorption of the antiserum with the peptide. Fig. 2l depicts a normal mammary artery with only minimal immunoreactivity of TLR-4 along the endothelial border.

Fig. 3 is not executed in color, and depicts the relative intensity of each band, at indicated dosage levels, of TLR-4 expression when analyzed by reverse transcription polymerase chain reaction ("RT-PCR"), relative to GAPDH expression in cultured human monocyte derived macrophages that were stimulated with either native or oxidized LDL for five hours.

DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENTS

Methods of the present invention inhibit Toll-like receptor-4 ("TLR-4") activity and expression by interfering with the production or biological activity of native TLR-4. One can use these methods to treat any disease in which inhibiting TLR-4 activity has a beneficial effect on a patient (e.g., ameliorating a disease, lessening the severity of its complications, preventing it from manifesting, preventing it from recurring, merely preventing it from worsening, or a therapeutic effort to affect any of the aforementioned, even if such therapeutic effort is ultimately unsuccessful). Diseases are known in the art in which TLR-4 activity is known or suspected to play a role in initiating, aggravating, or maintaining the pathological state that comprises the disease. Atherosclerosis, restenosis, inflammation and other vascular diseases are examples. Methods of the present invention may be used to treat any of these diseases.

In a preferred embodiment, methods of the present invention are used to inhibit atherosclerosis, transplant atherosclerosis, vein-graft atherosclerosis, stent restenosis, and angioplasty restenosis, and to thereby treat the cardiovascular diseases that atherosclerosis causes

(hereinafter "vascular diseases"). These methods may be used in any patient who could benefit from reducing atherosclerosis that is already present, from inhibiting atherosclerosis that has yet to form, or from both reducing existing atherosclerosis and inhibiting new atherosclerosis. Such patients include those suffering from, for example, angina pectoris and its subtypes (e.g., unstable
5 angina and variant angina); ischemias affecting organs such as the brain, heart, bone, and intestines, and conditions associated with the ischemias, such as stroke, transient ischemic attacks, heart attack, osteonecrosis, colitis, poor kidney function, and congestive heart failure; poor blood circulation to the extremities and the complications of poor blood circulation, such as slow wound healing, infections, and claudication; atherosclerosis itself, including restenosis following
10 angioplasty or stenting of atherosclerotic lesions; vein-graft atherosclerosis following bypass surgery; transplant atherosclerosis; and other diseases caused by or associated with atherosclerosis.

TLR-4 may be encoded by the RNA sequence set forth herein as SEQ ID NO. 6. Inhibition of this RNA, or those substantially similar to it, may correspondingly inhibit the
15 biological activity of TLR-4. Thus, various methods of the present invention are directed to inhibiting the expression of TLR-4 RNA.

The present invention contemplates a variety of TLR-4 inhibitors that are employed to inhibit the biological activity of TLR-4. These inhibitors may be administered to a mammal by any suitable means, such as those set forth in the various ensuing embodiments. Such inhibitors
20 may include any compound, pharmaceutical, or other composition that affects an inhibition of the biological activity of TLR-4. Such a composition may be administered to a mammal in an effective amount and by any suitable means, including, but not limited to, orally, topically, intravenously, intramuscularly, via a surgical device, such as a catheter, or via an implantable mechanism, such as a stent.

A first aspect of the present invention includes somatic cell gene transfer utilizing viral vectors containing TLR-4 gene sequences that express antisense RNA. Appropriate viral vectors that can express antisense TLR-4 RNA include expression vectors based on recombinant adenoviruses, adeno-associated viruses, retroviruses or lentiviruses, though non-viral vectors may be used, as well. An ideal vector for TLR-4 antisense gene transfer against atherosclerosis and angioplasty/stent-induced restenosis in mammals has the following attributes: (1) high efficacy of *in vivo* gene transfer; (2) recombinant gene expression in dividing as well as nondividing cells (the baseline mitotic rate in the coronary artery wall is <1% even in advanced lesions); (3) rapid and long-lived recombinant gene expression; (4) minimal vascular toxicity from inflammatory or immune responses; (5) absence of baseline immunity to the vector in the majority of the population; and (6) lack of pathogenicity of viral vectors. This is not to say that a vector must have all of these attributes; indeed, many useful vectors will not.

In a preferred embodiment of the invention, one employs adenovirus serotype 5 ("Ad5")-based vectors (available from Quantum Biotechnology, Inc., Montreal, Quebec, Canada) to deliver and express TLR-4 gene sequences expressing antisense RNA in cultured macrophages and vascular smooth muscle cells and in atherosclerosis-prone mice and swine. The recombinant Ad5 vectors have several advantages over other vectors such as liposomes and retroviruses. Unlike retroviral vectors, proliferation of the target cell is not required for infection by adenovirus vectors and thus, Ad5 vectors can infect cells *in vivo* in their quiescent state. Ad5 vectors are capable of infecting a number of different tissues although the transduction efficiency can vary according to the cell type. However, Ad5 vectors as a means of *in vivo* gene delivery have several drawbacks: (1) gene expression from cells transduced with the Ad5 vector is often transient due to the elimination of the Ad5-transduced cells by the host immune system; (2) Ad5 vectors may generate some toxicity to human recipients as observed in human clinical trials in cystic fibrosis patients; and (3) initial administration of Ad5 vectors produces blocking antibodies to the vectors,

thus repeated administrations of the adenoviral vector may not be effective. Even with these limitations, methods of the present invention utilize rAd5-mediated transfer of the TLR-4 sequence expressing antisense RNA. Using RT-PCR, a portion of TLR-4 is isolated and cloned upstream to the human cytomegalovirus ("CMV") major immediate early promoter-enhancer in a
5 direction to generate antisense TLR-4 RNA. The use of recombinant Ad5 vectors provides proof of the principle that adenovirus-mediated gene therapy might be particularly well suited as an adjunct to coronary angioplasty, since even temporary inhibition of smooth muscle cell proliferation might suffice to limit the formation of restenotic lesions.

A second aspect of the present invention provides a gene therapeutic method to produce
10 high levels of soluble forms of membrane bound TLR-4 that compete for non-bound HSP-60, LPS, and other ligands that are molecularly configured to operatively interact with a TLR-4 receptor, but lack at least a substantial portion of the TLR-4 signal transduction domain.

As with other types of disease, therapeutic strategies to treat atherosclerotic disease entail treatment for an extended period of time ranging from months to years. Prolonged and efficient
15 transgene transcription from heterologous promoters is a major consideration for gene therapies. The inclusion of a CMV promoter to drive expression of soluble TLR-4 in the present invention has been popularly used to express a variety of genes. It is, however, often subject to epigenetic silencing as are most promoters and transgenes. In an attempt to circumvent this problem, a variety of promoter expression strategies can be used to optimize the *in vivo* production of the
20 soluble TLR-4 in the present invention.

Efficient gene expression in viral vectors depends on a variety of factors. These include promoter strength, message stability and translational efficiency. Each of these factors must be explored independently to achieve optimal expression of a soluble TLR-4 gene. Applications of other promoter/enhancer variants to increase and optimize the expression of soluble TLR-4 *in*
25 *vitro* as well as *in vivo* are included within the scope of this invention. These include promoters

or enhancers stronger than CMV that exhibit inducibility such as tetracycline inducible promoters. Promoters/enhancers with tissue-specific functions that target, for example, vascular endothelial or smooth muscle tissue, and that produce sufficient amounts of soluble TLR-4 or its derivatives for a time and under condition sufficient to reduce the amount of TLR-4 ligand and thereby
5 inhibit the TLR-4 function may also be included. Levels and persistence of soluble TLR-4 expression can be compared with those obtained from the CMV promoter.

A third aspect of the present invention contemplates a somatic cell gene therapeutic method by administering a ribozyme-viral (adeno, adeno-associated or lentiviral) or non-viral vector against TLR-4 mRNA in a mammal, and in particular in humans for treating the conditions
10 referred to above. The method involves development of a hammerhead ribozyme expression cassette that targets a sequence of TLR-4 mRNA. Ribozymes are sequence-specific endoribonucleases that catalytically cleave specific RNA sequences, resulting in irreversible inactivation of the target mRNA, thereby inhibiting the gene expression. T. Cech, "Biological catalysis by RNA," *Ann Rev Biochem.* 55:599-629 (1986); J.J. Rossi, "Therapeutic ribozymes:
15 principles and applications," *Bio Drugs* 9:1-10 (1998). Ribozymes offer advantages over antisense ODN. For instance, ribozymes possess higher catalytic activity than ODN; a comparatively smaller quantity of ribozyme-containing active is thus required for inhibition of gene expression. Ribozymes can be delivered exogenously or can be expressed endogenously with the use of appropriate promoters in a viral vector. Methods of the present invention utilize a
20 hammerhead ribozyme directed to human TLR-4 mRNA. Desired quantity or the length of expression of the ribozyme-viral or non-viral vector can readily be determined by routine experimentation, as can the most effective and/or convenient route of administration.

In a fourth aspect of the present invention, there is provided a non-viral method to inhibit the expression of TLR-4. This method involves synthesis of pentadecamer ("15-mer") ODN
25 corresponding to the sense and antisense sequence of human TLR-4 mRNA. Pentadecamer ODN

are known to bind strongly to single-stranded regions of target mRNA. D. Jaskuski *et al.*, "Inhibition of cellular proliferation by antisense oligonucleotide to PCNA cyclin," *Science* 240:1544-1548 (1988). Such strong binding may correspondingly result in strong inhibition of the translation of mRNA.

5 In a preferred method of the present invention, ODN are synthesized on a nucleic acid synthesizer, such as the EXPIDITE Nucleic Acid Synthesizer (available from Applied Biosystems, Inc., Rockville, MD) and purified using standard protocols.

In a fifth aspect of the present invention, there is provided a method to inhibit the expression of TLR-4 by RNAi. This new approach to silencing a gene product by degrading a
10 corresponding RNA sequence is reportedly more effective than alternative gene silencing methodologies, including antisense and ribozyme-based strategies. The method involves the use of dsRNA that are sufficiently homologous to a portion of the TLR-4 gene product such that the dsRNA degrades mRNA that would otherwise affect the production of TLR-4. siRNA, a well-defined 21-base duplex RNA (obtained from Dharmacon Research, Inc., Boulder, CO), may
15 operate in conjunction with various cellular components to silence the TLR-4 gene product with sequence homology. RNAi is described in Hammond *et al.*, "Post-Transcriptional Gene-Silencing by Double-Stranded RNA," *Nature* 110-119 (2001); Sharp, P.A., "RNA interference - 2001," *Genes Dev.* 15:485-490 (2001); and Elbashir, *et al.*, "RNA interference is mediated by 21- and 22-nucleotide RNAs," *Genes Dev.* 15:188-200, each of which is incorporated by reference
20 herein in its entirety.

Efficient gene silencing may be achieved by employing siRNA duplexes which include sense and antisense strands each including approximately 21 nucleotides, and further paired such that they possess about a 19-nucleotide duplex region and about a 2-nucleotide overhang at each 3' terminus. Elbashir *et al.*, "Duplexes of 21-nucleotide RNAs mediate RNA interference in
25 cultured mammalian cells," *Nature* 411:494-498 (2001). It will be appreciated by one of skill in

the art of RNAi that alternately sized sense or antisense strands and/or variations on the size of the duplex and the overhang region that comprise them may be suitable for use with the methods of the present invention, and are contemplated as being within the scope thereof. Such appropriate alternate sizes may be readily ascertained without undue experimentation by one possessing such skill.

Furthermore, the inclusion of symmetric 3'-terminus overhangs may aid in the formation of specific endonuclease complexes ("siRNPs") with roughly equivalent ratios of sense and antisense target RNA cleaving siRNPs. It is believed that the antisense siRNA strand is responsible for target RNA recognition, while the 3'-overhang in the sense strand is not involved in this function. Therefore, in a preferred embodiment, the UU or dTdT 3'-overhang of an antisense sequence is complementary to target mRNA, however the symmetrical UU or dTdT 3'-overhang of the sense siRNA oligo need not correspond to the mRNA. Deoxythymidines may be included in either or both 3'-overhangs; this may increase nuclease resistance. However, siRNA duplexes that include either UU or dTdT overhangs may be equally resistant to nuclease.

The siRNA duplexes used in accordance with the present invention may be introduced to a cell via an appropriate viral or non-viral vector. Such vectors include those described above with regard to the somatic gene cell transfer embodiment of the present invention.

In a sixth aspect of the present invention, a method of inhibiting the TLR-4 cell-signaling pathway by peptide mimetics is provided. This method involves the introduction of small peptides (i.e., peptides of approximately 10-20 amino acids) that bind to TLR-4 ligands, thereby preventing these ligands from binding to a TLR-4 or associated receptor (e.g., MD2). TLR-4 is generally found on a cell surface substantially adjacent to an MD2 receptor. It is believed that, in order to initiate TLR-4 cell signal transduction, a TLR-4 ligand must bind simultaneously to the TLR-4 receptor as well as the adjacent MD2 receptor. Binding to only one of these receptors is insufficient to propagate TLR-4 cell signal transduction. Bulut *et al.*, "Chlamydial Heat Shock

Protein 60 Activates Macrophages and Endothelial Cells Through Toll-like Receptor 4 and MD2 in a MyD88-Dependent Pathway," *J. Immunol.*, 168:1435-1440.

The MD2 receptor is a protein constructed of approximately 133 individual amino acids. Short, overlapping segments (e.g., approximately 10-20 amino acids in length) of the MD2
5 receptor molecule may be separated to test which individual segments effect TLR-4 cell signal transduction by binding to a TLR-4 ligand. Segments are overlapping insofar as a portion of one end of one segment separated for testing corresponds to a portion of one end of a second segment separated for testing. Following separation, the segments are duplicated and tested to determine which comprise at least a portion of the MD2 receptor that binds to a TLR-4 ligand, such as
10 cHSP60, LPS, or other ligands that are molecularly configured to operatively interact with a TLR-4 receptor. A segment suitable for use in accordance with the method of the present invention comprises at least a portion of the MD2 receptor that binds to a TLR-4 ligand, such that the administration of a sufficient amount of individual copies of this segment will hinder TLR-4 signal transduction. Once administered, segments preferably bind to the MD2 binding sites of the
15 TLR-4 ligands, thereby preventing the ligands from binding to the corresponding sites on the MD2 receptor. This may significantly hinder TLR-4 cell signal transduction.

The same process may be implemented to identify a segment of the TLR-4 receptor that may similarly hinder TLR-4 signal transduction. Short, overlapping segments (e.g., approximately 10-20 amino acids in length) of the TLR-4 receptor molecule may be separated to
20 test which individual segments effect TLR-4 cell signal transduction by binding to a TLR-4 ligand. Following separation, the segments are duplicated and tested to determine whether the segment comprises at least a portion of the TLR-4 receptor that binds to a TLR-4 ligand, such as cHSP60, LPS, or other ligands that are molecularly configured to operatively interact with a TLR-4 receptor. A segment suitable for use in accordance with the method of the present invention
25 comprises at least a portion of the TLR-4 receptor that binds to a TLR-4 ligand, such that the

administration of a sufficient amount of individual copies of this segment will hinder TLR-4 signal transduction. Once administered, segments preferably bind to the binding sites of the TLR-4 ligands, thereby preventing the ligands from binding to the corresponding sites on the TLR-4 receptor. This may significantly hinder TLR-4 cell signal transduction.

5 In accordance with the method of the present invention, a segment that does, in fact, include at least a portion of the MD2 or TLR-4 receptor that binds to a TLR-4 ligand may be administered to a patient. Segments that include a portion of the MD2 receptor, segments that include a portion of the TLR-4 receptor, or combinations thereof may be administered. Moreover, administration may be performed by any suitable means, including via an oral form, such as a capsule, tablet, solution, or suspension; an intravenous form; an injectable form; an implantable
10 form, such as a stent coating, a sustained release mechanism, or a biodegradable polymer unit; or any other suitable mechanism by which an active or therapeutic agent may be delivered to a patient. The dosage may similarly be determined in accordance with the selected form of administration, the level of which may be readily ascertained without undue experimentation, as
15 can the most suitable means of administration.

In a seventh aspect of the present invention, a method of inhibiting TLR-4 expression through the introduction an anti-TLR-4 antibody is provided. Any suitable anti-TLR-4 antibody may be used in conjunction with this aspect of the present invention, including, but in no way limited to, anti-TLR-4 antibodies, and any suitable derivatives thereof, equivalents thereof, or
20 compounds with active sites that functions in a manner similar to anti-TLR-4 antibodies, whether those compounds are naturally occurring or synthetic (all hereinafter included within the term "anti-TLR-4 antibody").

An appropriate quantity of an anti-TLR-4 antibody necessary to affect the method of the present invention, and the most convenient route of delivering the same to a mammal may be
25 determined by one of ordinary skill in the art, without undue experimentation. Furthermore, it

will be readily appreciated by one of such skill that an anti-TLR-4 antibody may be formulated in a variety of pharmaceutical compositions, any one of which may be suitable for use in accordance with the method of the present invention.

Such an antibody may be delivered to a mammal through any conventional mechanism in
5 an amount effective to inhibit the TLR-4 signaling pathways in a mammal; the mechanism of delivery and quantity of antibody necessary for inhibiting TLR-4 expression both being readily ascertainable without undue experimentation.

The vascular delivery of TLR-4 inhibiting compositions composed in accordance with any of the various embodiments of the present invention can be accomplished by any of a wide range
10 of local delivery devices and methods. K. L. March, "Methods of local gene delivery to vascular tissues," *Semin Intervent Cardiol*, 1:215-223 (1996). Local delivery is preferred because, for those compositions that include a viral or non-viral vector, site-specific delivery may result in maximal therapeutic efficacy with minimal systemic side effects. These local delivery devices typically entail an endovascular or "inside-out" approach, whereby therapeutic agents are
15 delivered to the target site via intravascular catheters or devices. Although gene transfer is demonstrated for each device, most studies of catheter-based gene transfer reveal low efficiency, rapid redistribution of the infused material, and escape of the infusate into the systemic circulation.

Recently, several devices with modified needles capable of direct injection into interstitial
20 tissue of either myocardium or vasculature have been described. One such approach to local drug delivery is via the nipple balloon catheter, such as the INFILTRATOR® (available from InterVentional Technologies, Inc., San Diego, CA), although any appropriate catheter may be used. Methods of the present invention utilize the INFILTRATOR® for intramural delivery of small volumes of high-titer rAd5, where such a viral vector is appropriate. The INFILTRATOR®
25 catheter offers improved local gene delivery by placing vector particles directly and deeply within

the vascular wall. The INFILTRATOR® catheter is designed to provide direct intramural delivery of agents by mechanical access into the media and inner adventitia, which is achieved using sharp-edged injection orifices mounted on the balloon surface. P. Barath *et al.*, "Nipple balloon catheter," *Semin Intervent Cardiol*, 1:43 (1996). This catheter has been used clinically.

5 G. S. Pavlides *et al.*, "Intramural drug delivery by direct injection within the arterial wall: first clinical experience with a novel intracoronary delivery-infiltrator system," *Cathet Cardiovasc Diagn*, 41:287-292 (1997). Further, the INFILTRATOR® has been demonstrated to yield enhanced local transduction efficiency by adenoviral vectors compared with that which may be achieved by endoluminal delivery. T. Asahara *et al.*, "Local delivery of vascular endothelial

10 growth factor accelerates reendothelialization and attenuates intimal hyperplasia in balloon-injured rat carotid artery," *Circulation*, 91:2793-2801 (1995).

Methods of the present invention also utilize a perivascular or "outside-in" approach of drug delivery in the vessel wall by modifying the procedure applied in periadventitial carotid injury in a mouse, as described in Example 10 below. Oguchi S, *et al.* "Increased intimal

15 thickening after arterial injury in hypercholesterolemic apolipoprotein E-deficient mice: finding a novel method," *Circulation (supp.)* I-548:3066 (1997); P. Dimayuga *et al.*, "Reconstituted HDL containing human apolipoprotein A-1 reduces VCAM-1 expression and neointima formation following periadventitial cuff-induced carotid injury in apo E null mice" *Biochem Biophys Res Commun*. 264:465-468, (1999).

20 By directly targeting the genes involved via gene therapeutic approaches, methods of the present invention may be used in stent coatings that eliminate or substantially reduce restenosis following stent placement, as well as geometric miss and late in-stent thrombosis following intracoronary brachytherapy. Methods of the present invention contemplate stents coated with TLR-4 inhibiting compositions. As these gene therapeutic agents may be used as coatings on

25 already existing stents, they may be deployed without increasing procedure time, and will not

require significant additional equipment, expertise, hospitalization or expense. This strategy should prove cost-effective in the long run since, if successful, it will diminish the need for repeat hospitalizations and additional intervention procedures. Outcomes should also be favorable, to the extent that the strategy is effective in minimizing clinical events associated with restenosis following stent placement, and geometric miss and late in-stent thrombosis following intracoronary brachytherapy. Coated stents may eventually be implanted in all patients who are candidates for stents, since it is presently not possible to determine prior to the procedure which patients will suffer from restenosis or other complications associated with arterial injury following coronary intervention.

10 Since TLR-2 and TLR-4 play an important role in the innate immune and inflammatory response, we investigated the expression of these receptors, and found that TLR-4 exhibits preferential expression in lipid-rich and macrophage-infiltrated murine aortic and human coronary atherosclerotic plaques. The inventor's *in vitro* studies, described below, demonstrated basal expression of TLR-4 by macrophages, which was up-regulated by oxidized LDL ("ox-LDL").

15 While not wishing to be bound by any theory, these findings suggest a potential role for TLR-4 in lipid-mediated pro-inflammatory signaling in atherosclerosis. Moreover, as TLR-4 is a receptor that recognizes chlamydial antigens such as cLPS and cHSP-60, endotoxin, and other ligands that are molecularly configured to operatively interact with a TLR-4 receptor it may provide a molecular link between chronic infection, inflammation, and atherosclerosis.

20 The pro-inflammatory signaling receptor TLR-4 is expressed in lipid-rich, macrophage-infiltrated atherosclerotic lesions of mice and humans. Further, TLR-4 mRNA in cultured macrophages is up-regulated by ox-LDL but not native LDL ("N-LDL"). Together, these findings suggest that enhanced TLR-4 expression may play a role in inflammation in atherosclerosis.

Cells of the innate immune system, such as macrophages, have the ability to recognize common and conserved structural components of microbial origin by pattern recognition receptors. The human homologue of *Drosophila* Toll, TLR-4, is a pattern recognition receptor, which activates NF- κ B, and up-regulates a variety of inflammatory genes in response to microbial pathogens. Toll-like receptors play a fundamental role in the activation of innate immune responses and pathogen recognition. Further, activation of NF- κ B is essential for the regulation of a variety of genes involved in the inflammatory and proliferative responses of cells critical to atherogenesis. Both NF- κ B and genes regulated by NF- κ B are expressed in atherosclerotic lesions. Since NF- κ B activation leads to transcription of a number of pro-inflammatory genes involved in athero-thrombosis, it may be that infectious agents and clamydial antigens such as LPS and/or HSP-60 contribute to enhanced and chronic inflammation by signaling through the TLR-4 receptor, which is up-regulated by ox-LDL.

The inventor's findings of increased expression of TLR-4 induced by ox-LDL suggests a potential mechanism for the synergistic effects of hypercholesterolemia and infection in acceleration of atherosclerosis observed in experimental models and human epidemiologic observations. This provides new insight into the link among lipids, infection/inflammation and atherosclerosis.

EXAMPLE 1

Preparation of Mouse Tissue

Five apolipoprotein E-deficient ("apoE -/-") mice (C57BL/6J strain, aged 5 weeks, 18 to 20 grams; obtained from Jackson Laboratory, Bar Harbor, ME) were fed a high fat, high cholesterol (i.e., atherogenic) diet containing 42% (wt/wt) fat and 0.15% cholesterol from 6 weeks of age through the duration of the experiment. After anesthesia with ETHRANE (available from Abbot Laboratories, Abbott Park, IL), the mice were sacrificed at 26 weeks of age, and their

hearts and proximal aortas (including ascending aorta, aortic arch and a portion of descending aorta) were excised and washed in phosphate-buffered saline ("PBS") to remove blood. The basal portion of the heart and proximal aorta were embedded in OCT compound using TISSUE-TEK VIP (available from Sakura Finetek USA, Inc., Torrance, CA), frozen on dry ice and then stored
5 at -70°C until sectioning. Serial 10µm-thick cryosections (every fifth section from the lower portion of the ventricles to the appearance of aortic valves, every other section in the region of the aortic sinus, and every fifth section from the disappearance of the aortic valves to the aortic arch) were collected on poly-D-lysine-coated slides (available from Becton Dickinson & Co., Franklin Lakes, NJ). Sections were stained with Oil Red O and hematoxylin, and counterstained with Fast
10 Green (all available from Sigma Chemical Co., St. Louis, MO, "Sigma") for the identification of atheromatous lesions, arterial wall calcification, and cartilaginous metaplasia. The presence of calcium deposits was confirmed by the alizarin red S (available from Sigma) and von Kossa techniques using representative sections.

15

EXAMPLE 2

Preparation of Human Tissue and Human Monocyte-Derived Macrophages

Human coronary artery specimens from nine autopsy cases were collected within 24 hours of death, fixed with 10% formalin (available from Sigma) overnight and embedded in paraffin. Five of the nine coronary artery specimens included lipid-rich plaques containing a well-defined
20 lipid-core covered by a fibrous cap, and the other four of the nine specimens included fibrous plaques, which contained mostly extracellular matrix without a lipid-core. Normal mammary artery specimens were also obtained from four additional autopsy cases. Five µm-thick sections were cut and applied to slides for both hematoxyline-eosin and immunohistochemical staining. Peripheral blood monocytes were isolated from whole blood of normal human subject by
25 FICOLL-PAQUE density gradient centrifugation (available from Pharmacia LKB Biotechnology,

Inc., Piscataway, NJ). Monocyte-derived macrophages were cultured in RPMI 1640 (available from Sigma) containing 10% fetal calf serum ("FCS"), 100 U/ml penicillin, 100 µg/ml streptomycin and 0.25 µg/ml amphotericin B for 5 days and then starved in the culture medium without FCS but with 0.1% low endotoxin bovine serum albumin ("BSA") (obtained from Sigma).

EXAMPLE 3

Immunohistochemistry

Frozen sections of the apoE ^{-/-} mouse aortic root were fixed with acetone for 5 minutes at room temperature and then immunostained with rabbit anti-hTLR-4 immune serum (1:100; obtained from Ruslan Medzhitov, Asst. Prof. of Immunobiology, Yale University, New Haven, CT) following the instructions on the immunostaining kit available from DAKO (Carpinteria, CA, "DAKO"). Rat anti-mouse macros Ab (1:500; available from Serotec, U.K.) were used as macrophage marker. Colors were developed using the DAKO AES substrate system. Smooth muscle cells were stained by a mouse anti-actin Ab conjugated with alkaline phosphatase (1:50, available from Sigma). Colors were developed using VECTOR Red Alkaline Phosphatase Substrate Kit I (obtained from Vector Laboratories, Inc., Burlingame, CA). Rabbit IgG or rabbit serum was used as a negative control.

For human atherosclerotic plaques, following deparaffinization in graded alcohol, sections were immunostained using rabbit anti-human TLR-4 and TLR-2 antiserum (1:100) raised against extracellular peptide domains of TLR-4 and TLR-2 (available from Berkeley Antibody Company, Richmond, CA). Following immunoperoxidase staining, the representative fields were photographed. Cells were lysed in Laemmli buffer and separated with a 10% sodium dodecyl sulphate polyacrylamide gel electrophoresis ("SDS-PAGE"). The protein was then transferred onto a polyvinylidene difluoride membrane, and the membrane was probed with anti-TLR-2, anti-

TLR-4 antibodies, and prebleeds corresponding to each antibody (1:2,000). After incubation with horseradish peroxidase-conjugated goat anti-rabbit antibody (available from Rockland Immunochemicals for Research, Gilbertsville, PA), the membrane was developed with an enhanced chemiluminescence ECL Western Blotting Detection Kit (available from Amersham Pharmacia Biotech UK Ltd., Buckinghamshire, England). Pre-incubating the anti-TLR-4 serum with TLR-4 peptide (SEQ ID NO. 5) was used to demonstrate specificity of the strain and rabbit IgG or rabbit serum instead of primary antibody was used as a negative control.

EXAMPLE 4

10

Double Immunohistochemistry

Double immunostaining of human atherosclerotic plaques was performed using an EnVision Doublestain System (available from DAKO). Following TLR-4 immunostaining, 3,3'-diaminobenzadine (obtained from Sigma) was used as the peroxidase chromogenic substrate. Mouse monoclonal anti-human CD68 antibody (360 µg/ml, 1:20 dilution; available from DAKO) for macrophages and mouse monoclonal anti-human α-actin antibody (100 µg/ml, 1:100 dilution; available from DAKO) for smooth muscle cells were used with Fast Red (available from Sigma) as the alkaline phosphatase chromogenic substrate.

EXAMPLE 5

20

Preparation and Modification of Lipoproteins

Human N-LDL (obtained from Sigma) was dialyzed against isotonic phosphate saline buffer (pH 7.4) to remove ethylenediamine tetraacetic acid ("EDTA") by using a 10,000 molecular weight cut-off SLIDE-A-LYZER dialysis cassette (obtained from Pierce Chemical Co., Rockford, IL). Ox-LDL was prepared by incubating 0.1 mg of LDL protein/ml with 5 µM of copper sulfate (CuSO₄) for 24 hours at 37°C, and stopped by adding butylated hydroxytoluene

(2,6-di-*t*-butyl-*p*-cresol) (available from Sigma) to a final concentration of 0.1mM. Ox-LDL was separated from CuSO₄ and equilibrated into the cell culture medium over a PD-10 column (available from Pharmacia Fine Chemicals, Uppsala, Sweden). All reagents were endotoxin-free. LPS levels of LDL preparations were confirmed with a chromogenic *Limulus* assay and contained
5 less than 0.3 pg of LPS/μg of LDL protein.

The extent of oxidation of the lipoprotein preparations was determined by a thiobarbituric acid reactive substance ("TBARS") assay. Concentrated trichloroacetic acid was added to aliquots of lipoprotein samples containing 1.5 mg of protein to give a final concentration of 5%. An equal volume of 1% thiobarbituric acid was then added and the mixture was heated in a water
10 bath at 100°C for 20 min. After centrifugation to clarify the solution, the peak absorbance at 582 nm was read on a Beckman DB Spectrophotometer (available from Beckman Coulter, Inc., Fullerton, CA) against a buffer blank. The amount of thiobarbituric-reactive substance was calculated from a standard curve, with malonaldehyde bis(dimethylacetal) (available from Sigma) as the standard. The ox-LDL had 20-25 nM TBARS/mg of cholesterol.

15

EXAMPLE 6

Reverse Transcription-Polymerase Chain Reaction (RT-PCR)

Total RNA was isolated from resting N-LDL, ox-LDL stimulated human monocyte-derived macrophage cells using an RNA Stat60 isolation reagent (obtained from Tel-test 'B', Inc.,
20 Friendswood, TX) following manufacturer's instruction and treated with RNase-free DNase I. For RT reaction, the SUPERScript MMLV preamplification system (obtained from Life Technologies, Inc., Gaithersburg, MD) was applied. PCR amplification was performed with TAQ GOLD polymerase (obtained from Perkin Elmer, Foster City, CA) for 32 cycles at 95°C for 45s, 54°C for 45s, and 72°C for 60s (for TLR-2 and TLR-4). The oligonucleotide primers used for RT-
25 PCR for TLR-2 were SEQ ID NO. 1 and SEQ ID NO. 2, and for TLR-4 were SEQ ID NO. 3 and

SEQ ID NO. 4. Glyceraldehyde-3-phosphate dehydrogenase ("GAPDH") primers were obtained from Clontech Laboratories, Inc. (Palo Alto, CA).

The TLR-2 and TLR-4 RT-PCR fragments were purified and sequenced to confirm the identity of the fragments. Real-time quantitative PCR was performed on an iCycler Thermal
5 Cycler (obtained from Bio-Rad Laboratories, Inc., Hercules, CA) using an SYBR Green RT-PCR Reagents kit (obtained from Applied Biosystems, Foster City, CA) and the TLR primers described above. The semi-quantitative RT-PCR experiments were repeated with cells pretreated for 1 hour with 15d-PGJ₂ (20 μM), proteasome inhibitor I (100 μM) (available from Affinity Bioreagents, Inc., Golden, CO), or cycloheximide (10 μm/ml). Endothelial cells were pretreated with NF-κB
10 p65 antisense and sense oligonucleotides (30 μM) for 24-48 hours, three times before LPS stimulation (50 ng/ml). For densitometry analysis, the intensity of the bands were measured by Digital Science 1D Image Analysis Software (obtained from Eastman Kodak Co., Rochester, NY) and normalized with GAPDH intensity.

15

EXAMPLE 7

TLR-4 is Expressed in Atherosclerotic Lesions of the ApoE -/- Mice

As depicted in Fig. 1, all five apoE -/- mice exhibited TLR-4 immunoreactivity in the atherosclerotic lesions of the aortic root, which co-localized with macrophage immunoreactivity. TLR-4 staining was absent in the normal vessels obtained from control C56BL/6J mice (Fig. 1e).
20 Mouse IgG staining was negative and pre-incubation of the tissue sections with the specific peptide against which the anti-TLR-4 antiserum was generated completely blocked the TLR-4 staining in the apoE -/- vessels, indicating the specific nature of the TLR-4 immunostaining. No TLR-2 immunoreactivity was observed in normal or atherosclerotic lesions (not shown).

EXAMPLE 8

TLR-4 is Expressed in Human Coronary Plaques

The human coronary atherosclerotic plaques were classified into lipid-rich plaques containing a well-defined lipid-core covered by a fibrous cap (n=5), and fibrous plaques which contained mostly extracellular matrix without a lipid-core (n=4). As depicted in Fig. 2, strong TLR-4 expression (brown staining) was observed around the lipid core at the shoulder of lipid-rich plaques where it co-localized with macrophage immunoreactivity. Incubation of the antiserum with the peptide used to generate the primary antibody blocked TLR-4 immunoreactivity, confirming the specificity of the anti-TLR-4 antiserum. Double staining showed close spatial co-localization of TLR-4 expression with macrophage immunoreactivity. No TLR-4 immunoreactivity or macrophage immunoreactivity was found in fibrous plaques, which demonstrated strong smooth muscle α -actin immunoreactivity. Normal mammary arteries showed only minimal or no TLR-4 expression. TLR-2 immunoreactivity was absent in all plaques while control staining was positive in THP-1 cells (not shown).

EXAMPLE 9

TLR-4 mRNA Regulation by Ox-LDL

Cultured human monocyte derived macrophages were stimulated with N-LDL or ox-LDL for 5 hours. RT-PCR was performed for TLR-2 and TLR-4, and relative intensity was calculated by densitometry as described in Faure *et al.*, at 2018-2024. As depicted in Fig. 3, RT-PCR showed basal TLR-2 and TLR-4 mRNA expression by macrophages. The TLR-4 mRNA was upregulated by ox-LDL in a dose-dependent manner and up to threefold, whereas N-LDL had no effect. TLR-2 mRNA was not upregulated by ox-LDL.

EXAMPLE 10Perivascular or "Outside In" Approach to Drug Delivery

ApoE -/- mice (20 weeks of age, 6 per group) were anesthetized, and the carotid artery was exposed by making a small incision in the side of the neck. A section of artery was loosely
5 sheathed with a cuff made of a TYGON tube (3.0 mm long, 0.5 mm inner diameter; obtained from Saint-Gobain Performance Plastics, Wayne, NJ). A biodegradable biocompatible polymeric material, ATRIGEL (obtained from Atrix Laboratories, Ft. Collins, CO), a copolymer of polylactic and polyglycolic acid, was used for the local delivery of viral particles. An 18% (w/w) polymeric gel in PBS with 1×10^8 pfu of rAd5 (right carotid) or without rAd5 (left carotid) was
10 applied between the cuff and the vessel using a syringe and blunt cannula. The gel compound used in the study was a free-flowing liquid below body temperature. When placed in an aqueous environment at or above body temperature, the viscosity increases and the gel solidifies into a viscous mass. Once applied to the artery *in vivo*, the polymer turns into a gel immediately on contact and the gel is gradually resorbed in about 14 to 21 days, thereby providing potential use as
15 a drug depot.

WHAT IS CLAIMED IS:

1. A system for inhibiting the biological activity of Toll-like receptor-4 (TLR-4) comprising:

an intravascular device; and

- 5 a therapeutic composition coated upon the intravascular device, the therapeutic composition comprising a TLR-4 inhibitor.

2. The system of claim 1, wherein the intravascular device is selected from the group consisting of a catheter and a stent.

3. The system of claim 1, wherein the TLR-4 inhibitor is selected from the group
10 consisting of a nucleic acid expressing antisense TLR-4 RNA, a nucleic acid encoding a soluble TLR-4 protein, a nucleic acid encoding a hammerhead ribozyme that cleaves TLR-4 mRNA, an antisense TLR-4 oligodeoxynucleotide (ODN), a nucleic acid expressing a double stranded RNA (dsRNA) that is sufficiently homologous to a portion of a TLR-4 gene product such that the dsRNA is capable of inhibiting the encoding function of mRNA
15 that would otherwise cause the production of TLR-4, a protein sequence that corresponds to at least a portion of a receptor that binds to a TLR-4 ligand during a TLR-4 signal transduction event, and an anti-TLR-4 antibody.

4. The system of claim 3, wherein the TLR-4 inhibitor is the nucleic acid expressing antisense TLR-4 RNA.

- 20 5. The system of claim 3, wherein the TLR-4 inhibitor is the nucleic acid encoding the hammerhead ribozyme that cleaves TLR-4 mRNA.

6. The system of claim 3, wherein the TLR-4 inhibitor is the antisense TLR-4 oligodeoxynucleotide (ODN).
7. The system of claim 3, wherein the TLR-4 inhibitor is the anti-TLR-4 antibody.
8. The system of claim 1, wherein the TLR-4 inhibitor is included within a vector.
- 5 9. The system of claim 8, wherein the vector is selected from the group consisting of adenoviruses, adeno-associated viruses, retroviruses, lentiviruses, viral vectors, and non-viral vectors.
- 10 10. The system of claim 8, wherein the vector is an adenovirus serotype 5-based vector.
11. The system of claim 8, wherein the TLR-4 inhibitor is selected from the group consisting of a nucleic acid expressing antisense TLR-4 RNA, a nucleic acid encoding soluble TLR-4 protein, a nucleic acid encoding a hammerhead ribozyme that cleaves TLR-4 mRNA, and a nucleic acid expressing a double stranded RNA (dsRNA) that is sufficiently homologous to a portion of a TLR-4 gene product such that the dsRNA is capable of inhibiting the encoding function of mRNA that would otherwise cause the production of TLR-4.
- 15 12. The system of claim 1, further comprising an amount of the therapeutic composition sufficient to inhibit a vascular disease.
13. The system of claim 12, wherein the vascular disease is selected from the group consisting of atherosclerosis, transplant atherosclerosis, vein-graft atherosclerosis, thrombosis, restenosis, stent restenosis, and angioplasty restenosis.
- 20

14. The system of claim 3, wherein the TLR-4 inhibitor is the nucleic acid encoding the soluble TLR-4 protein.
15. The system of claim 14, wherein the soluble TLR-4 protein is unable to participate in normal TLR-4 signal transduction.
- 5 16. The system of claim 14, wherein the soluble TLR-4 protein lacks a substantial portion of the normal TLR-4 signal transduction domain.
17. The system of claim 14, wherein the soluble TLR-4 protein competes for a non-bound TLR-4 ligand.
18. The system of claim 17, wherein the non-bound TLR-4 ligand is a chlamydial heat
10 shock protein-60 (cHSP60) or a lipopolysaccharide (LPS).
19. The system of claim 3, wherein the TLR-4 inhibitor is the nucleic acid expressing the dsRNA, and the dsRNA further includes:
- a sense strand further including approximately 21 nucleotides; and
- an antisense strand further including approximately 21 nucleotides.
- 15 20. The system of claim 19, wherein the sense strand and the antisense strand are paired such that they possess a duplex region of approximately 19 nucleotides.
21. The system of claim 19, wherein the sense strand and the antisense strand each further include an overhang at a 3'-terminus of approximately 2 nucleotides.
22. The system of claim 21, wherein the sense overhang and the antisense overhang
20 are symmetrical.

23. The system of claim 21, wherein the antisense overhang comprises a UU 3'-overhang or a dTdT 3'-overhang.
24. The system of claim 23, wherein the UU 3'-overhang or the dTdT 3'-overhang is complementary to the mRNA.
- 5 25. The system of claim 21, wherein at least one of the sense overhang and the antisense overhang further includes a deoxythymidine.
26. The system of claim 3, wherein the TLR-4 inhibitor is the protein sequence that corresponds to at least the portion of the receptor that binds to the TLR-4 ligand during the TLR-4 signal transduction event.
- 10 27. The system of claim 26, wherein the receptor is a TLR-4 receptor or an MD2 receptor.
28. The system of claim 26, wherein the protein sequence comprises from about 10 to about 20 amino acids.
29. A method of treating a vascular disease, the method comprising the steps of:
- 15 providing a TLR-4 inhibitor; and
- administering the TLR-4 inhibitor to a mammal in an amount effective to at least partially inhibit the biological activity of TLR-4.
30. The method of claim 29, wherein the vascular disease is selected from the group consisting of atherosclerosis, transplant atherosclerosis, vein-graft atherosclerosis, thrombosis, restenosis, stent restenosis, and angioplasty restenosis.
- 20

31. The method of claim 29, wherein the step of administering the TLR-4 inhibitor further comprises administering the TLR-4 inhibitor in an amount effective to inhibit the vascular disease.

32. The method of claim 29, wherein the step of administering the TLR-4 inhibitor further comprises administering the TLR-4 inhibitor intravenously.

33. The method of claim 29, wherein the step of administering the TLR-4 inhibitor further comprises administering the TLR-4 inhibitor intramuscularly.

34. The method of claim 29, wherein the step of administering the TLR-4 inhibitor further comprises delivering the TLR-4 inhibitor with an intravascular device.

35. The method of claim 34, wherein the intravascular device is a catheter or a stent.

36. The method of claim 34, wherein the intravascular device is coated with the TLR-4 inhibitor.

37. The method of claim 29, wherein the TLR-4 inhibitor is selected from the group consisting of a nucleic acid expressing antisense TLR-4 RNA, a nucleic acid encoding a soluble TLR-4 protein, a nucleic acid encoding a hammerhead ribozyme that cleaves TLR-4 mRNA, an antisense TLR-4 oligodeoxynucleotide (ODN), a nucleic acid expressing a double stranded RNA (dsRNA) that is sufficiently homologous to a portion of a TLR-4 gene product such that the dsRNA is capable of inhibiting the encoding function of mRNA that would otherwise cause the production of TLR-4, a protein sequence that corresponds to at least a portion of a receptor that binds to a TLR-4 ligand during a TLR-4 signal transduction event, and an anti-TLR-4 antibody.

38. The method of claim 37, wherein the TLR-4 inhibitor is the nucleic acid expressing antisense TLR-4 RNA.

39. The method of claim 37, wherein the TLR-4 inhibitor is the nucleic acid encoding the hammerhead ribozyme that cleaves TLR-4 mRNA.

5 40. The method of claim 37, wherein the TLR-4 inhibitor is the antisense TLR-4 oligodeoxynucleotide (ODN).

41. The method of claim 37, wherein the TLR-4 inhibitor is the anti-TLR-4 antibody.

42. The method of claim 37, wherein the TLR-4 inhibitor is included within a vector.

10 43. The method of claim 42, wherein the vector is selected from the group consisting of adenoviruses, adeno-associated viruses, retroviruses, lentiviruses, viral vectors, and non-viral vectors.

44. The method of claim 42, wherein the vector is an adenovirus serotype 5-based vector.

15 45. The method of claim 42, wherein the TLR-4 inhibitor is selected from the group consisting of a nucleic acid expressing antisense TLR-4 RNA, a nucleic acid encoding soluble TLR-4 protein, a nucleic acid encoding a hammerhead ribozyme that cleaves TLR-4 mRNA, and a nucleic acid expressing a double stranded RNA (dsRNA) that is sufficiently homologous to a portion of a TLR-4 gene product such that the dsRNA is capable of inhibiting the encoding function of mRNA that would otherwise cause the
20 production of TLR-4.

46. The method of claim 37, wherein the TLR-4 inhibitor is the nucleic acid encoding the soluble TLR-4 protein.

47. The method of claim 46, wherein the soluble TLR-4 protein is unable to participate in normal TLR-4 signal transduction.

5 48. The method of claim 46, wherein the soluble TLR-4 protein lacks a substantial portion of the normal TLR-4 signal transduction domain.

49. The method of claim 46, wherein the soluble TLR-4 protein competes for a non-bound TLR-4 ligand.

10 50. The method of claim 49, wherein the non-bound TLR-4 ligand is a chlamydial heat shock protein-60 (cHSP60) or a lipopolysaccharide (LPS).

51. The method of claim 37, wherein the TLR-4 inhibitor is the nucleic acid expressing the dsRNA, and the dsRNA further includes:

a sense strand further including approximately 21 nucleotides; and

an antisense strand further including approximately 21 nucleotides.

15 52. The method of claim 51, wherein the sense strand and the antisense strand are paired such that they possess a duplex region of approximately 19 nucleotides.

53. The method of claim 52, wherein the sense strand and the antisense strand each further include an overhang at a 3'-terminus of approximately 2 nucleotides.

20 54. The method of claim 53, wherein the sense overhang and the antisense overhang are symmetrical.

55. The method of claim 53, wherein the antisense overhang comprises a UU 3'-overhang or a dTdT 3'-overhang.

56. The method of claim 55, wherein the UU 3'-overhang or the dTdT 3'-overhang is complementary to the mRNA.

5 57. The method of claim 53, wherein at least one of the sense overhang and the antisense overhang further includes a deoxythymidine.

58. The method of claim 37, wherein the TLR-4 inhibitor is the protein sequence that corresponds to at least the portion of the receptor that binds to the TLR-4 ligand during the TLR-4 signal transduction event.

10 59. The method of claim 58, wherein the receptor is a TLR-4 receptor or an MD2 receptor.

60. The method of claim 58, wherein the protein sequence comprises from about 10 to about 20 amino acids.

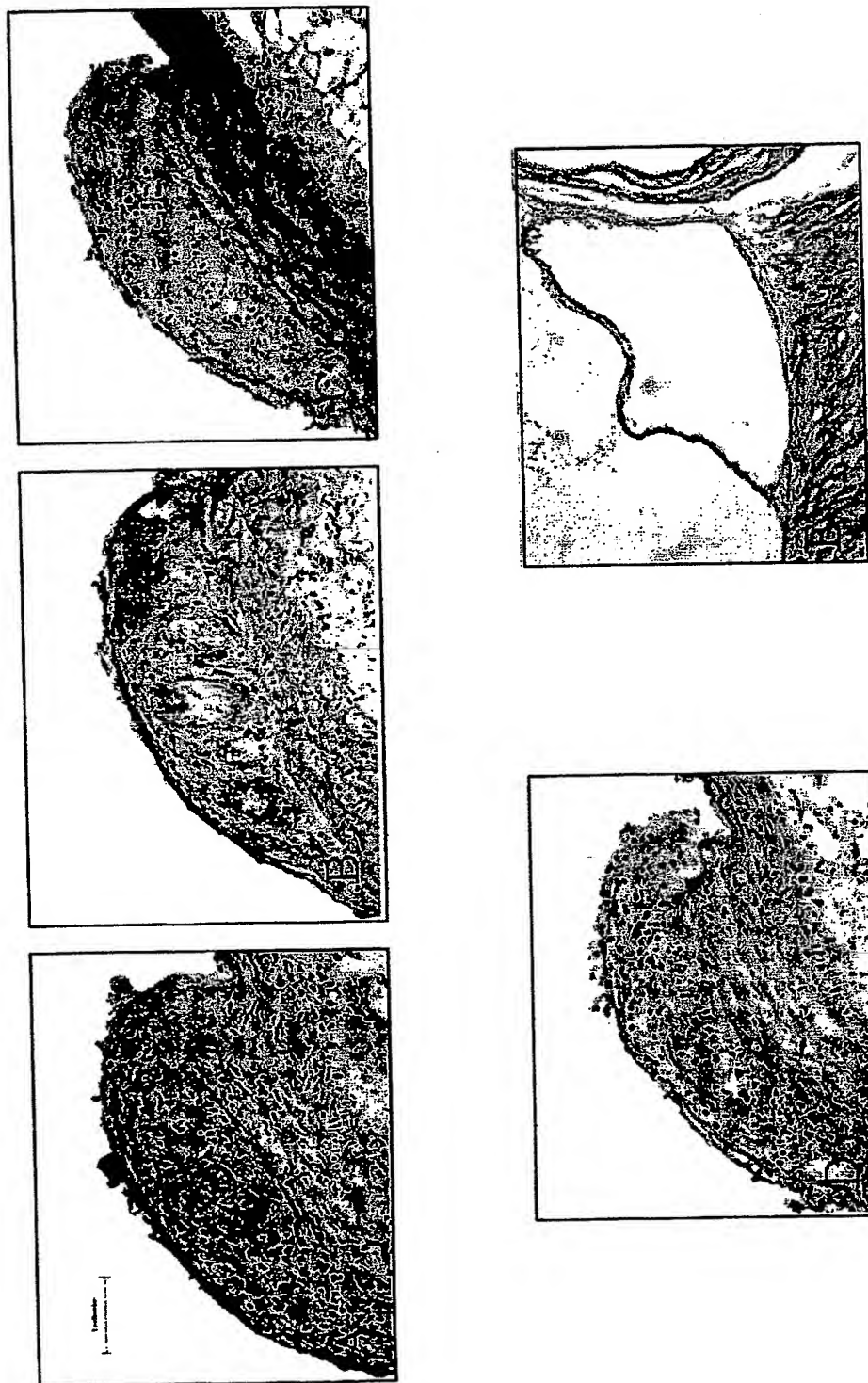


FIG. 1

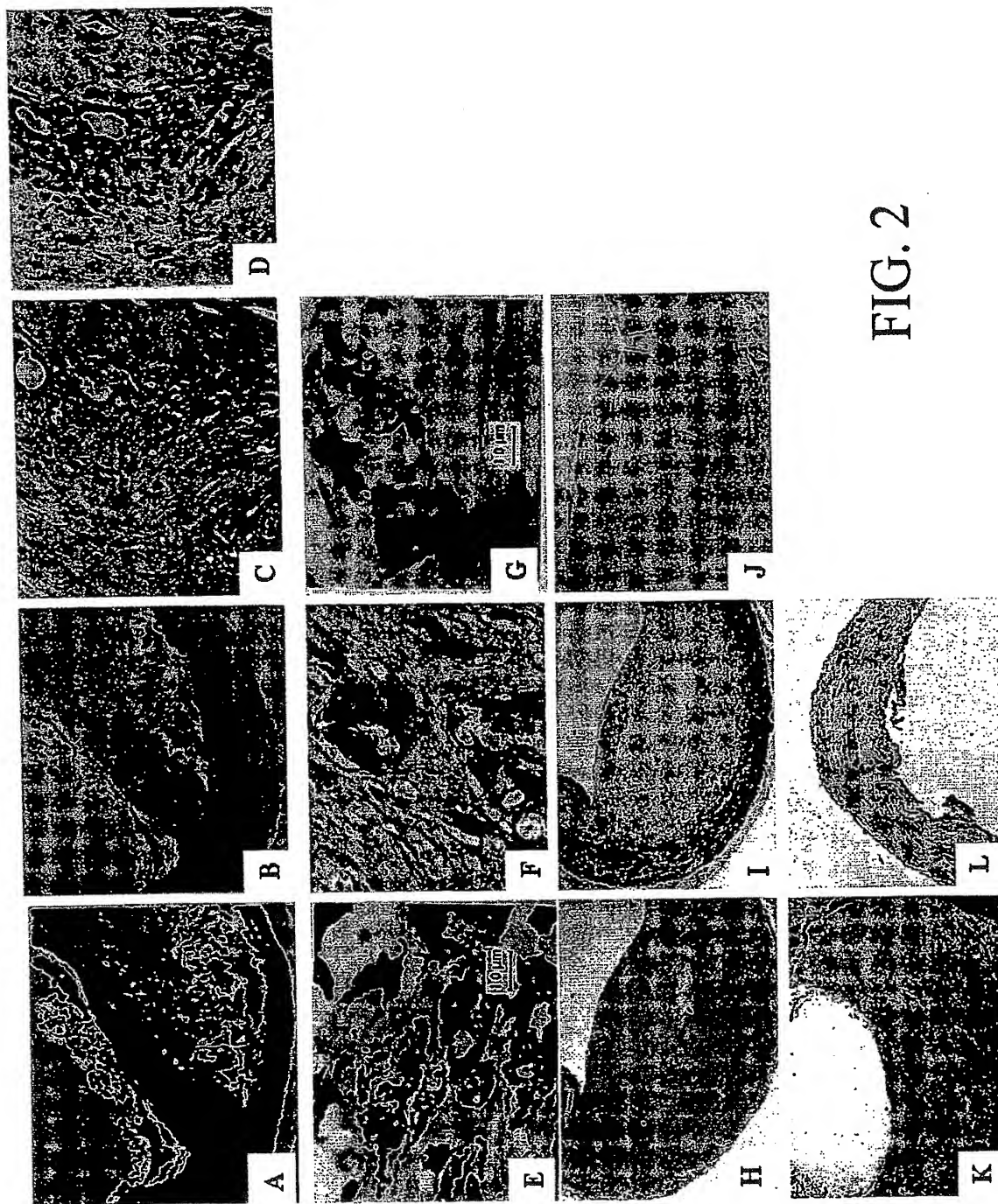
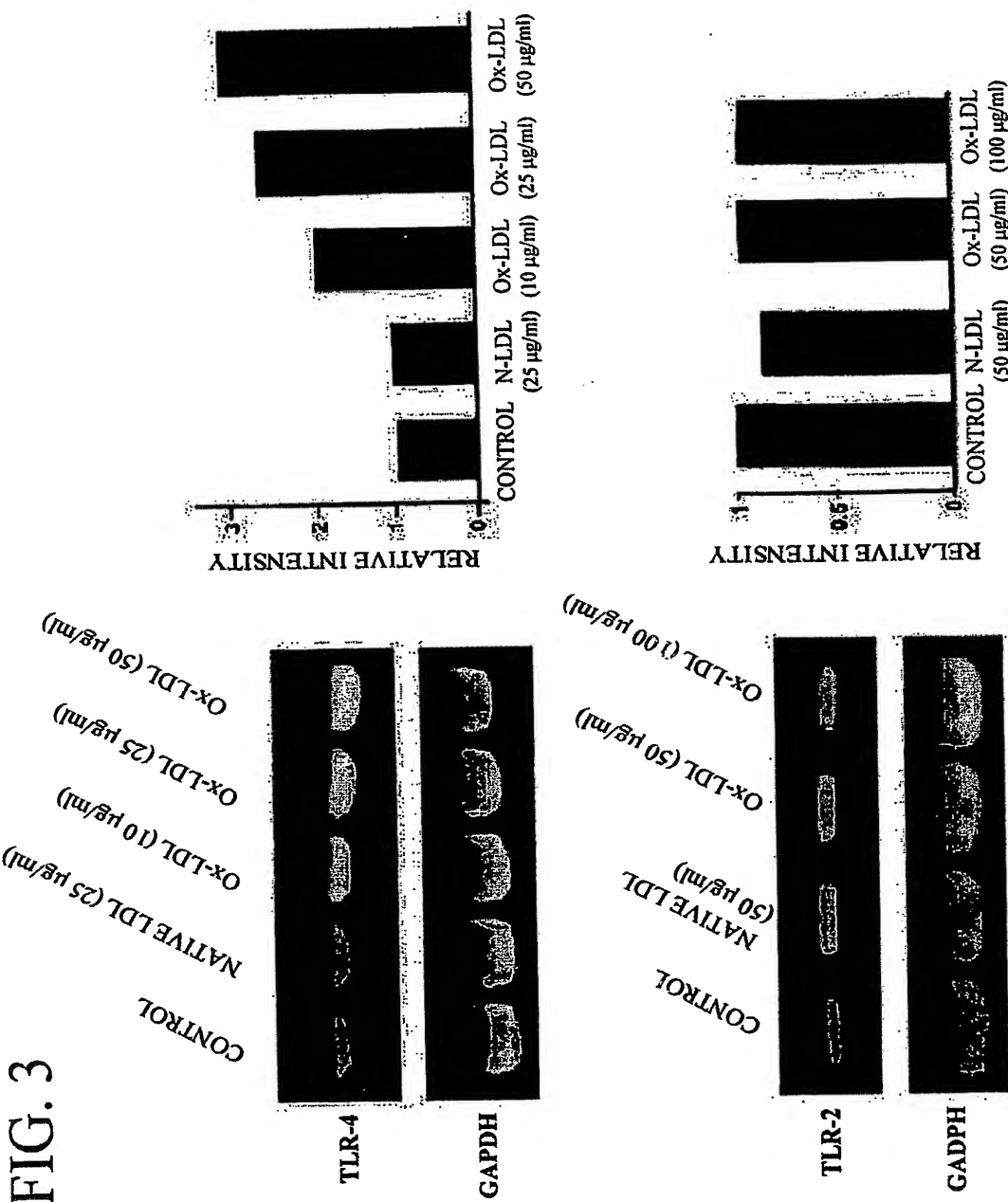


FIG. 2

FIG. 3



Sequence Data File for PCT.app.txt
SEQUENCE LISTING

<110> ARDITI, MOSHE
RAJAVASHISTH, TRIPATHI
SHAH, PREDIMAN K.

<120> METHODS FOR TREATING VASCULAR DISEASE BY INHIBITING
TOLL-LIKE RECEPTOR-4

<130> 81476-0140483

<140>

<141>

<160> 7

<170> PatentIn Ver. 2.1

<210> 1

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 1

gccaaagtct tgattgattg g

21

<210> 2

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 2

ttgaagttct ccagctcctg

20

<210> 3

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 3

tggtatcggt tccttataag

20

<210> 4

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 4

Sequence Data File for PCT.app.txt

19

gaaatggagg cacccttc

<210> 5
 <211> 23
 <212> PRT
 <213> Unknown Organism

<220>
 <223> Description of Unknown Organism: TLR-4 peptide

<400> 5
 Phe Lys Glu Ile Arg His Lys Leu Thr Leu Arg Asn Asn Phe Asp Leu
 1 5 10 15
 Ser Leu Asn Val Met Lys Thr
 20

<210> 6
 <211> 2400
 <212> DNA
 <213> Homo sapiens

<220>
 <221> CDS
 <222> (1)..(2397)

<400> 6
 atg gag ctg aat ttc tac aaa atc ccc gac aac ctc ccc ttc tca acc 48
 Met Glu Leu Asn Phe Tyr Lys Ile Pro Asp Asn Leu Pro Phe Ser Thr
 1 5 10 15
 aag aac ctg gac ctg agc ttt aat ccc ctg agg cat tta ggc agc tat 96
 Lys Asn Leu Asp Leu Ser Phe Asn Pro Leu Arg His Leu Gly Ser Tyr
 20 25 30
 agc ttc ttc agt ttc cca gaa ctg cag gtg ctg gat tta tcc agg tgt 144
 Ser Phe Phe Ser Phe Pro Glu Leu Gln Val Leu Asp Leu Ser Arg Cys
 35 40 45
 gaa atc cag aca att gaa gat ggg gca tat cag agc cta agc cac ctc 192
 Glu Ile Gln Thr Ile Glu Asp Gly Ala Tyr Gln Ser Leu Ser His Leu
 50 55 60
 tct acc tta ata ttg aca gga aac ccc atc cag agt tta gcc ctg gga 240
 Ser Thr Leu Ile Leu Thr Gly Asn Pro Ile Gln Ser Leu Ala Leu Gly
 65 70 75 80
 gcc ttt tct gga cta tca agt tta cag aag ctg gtg gct gtg gag aca 288
 Ala Phe Ser Gly Leu Ser Ser Leu Gln Lys Leu Val Ala Val Glu Thr
 85 90 95
 aat cta gca tct cta gag aac ttc ccc att gga cat ctc aaa act ttg 336
 Asn Leu Ala Ser Leu Glu Asn Phe Pro Ile Gly His Leu Lys Thr Leu
 100 105 110
 aaa gaa ctt aat gtg gct cac aat ctt atc caa tct ttc aaa tta cct 384
 Lys Glu Leu Asn Val Ala His Asn Leu Ile Gln Ser Phe Lys Leu Pro
 115 120 125
 gag tat ttt tct aat ctg acc aat cta gag cac ttg gac ctt tcc agc 432
 Glu Tyr Phe Ser Asn Leu Thr Asn Leu Glu His Leu Asp Leu Ser Ser

Page 2

Sequence Data File for PCT.app.txt

130	aac aag att caa agt att tat tgc aca gac ttg cgg gtt cta cat caa	480
Asn Lys Ile Gln Ser Ile Tyr Cys Thr Asp Leu Arg Val Leu His Gln		
145	150	155
160		
atg ccc cta ctc aat ctc tct tta gac ctg tcc ctg aac cct atg aac	528	
Met Pro Leu Leu Asn Leu Ser Leu Asp Leu Ser Leu Asn Pro Met Asn		
165	170	175
180		
ttt atc caa cca ggt gca ttt aaa gaa att agg ctt cat aag ctg act	576	
Phe Ile Gln Pro Gly Ala Phe Lys Glu Ile Arg Leu His Lys Leu Thr		
185	190	
195		
tta aga aat aat ttt gat agt tta aat gta atg aaa act tgt att caa	624	
Leu Arg Asn Asn Phe Asp Ser Leu Asn Val Met Lys Thr Cys Ile Gln		
200	205	
210		
ggt ctg gct ggt tta gaa gtc cat cgt ttg gtt ctg gga gaa ttt aga	672	
Gly Leu Ala Gly Leu Glu Val His Arg Leu Val Leu Gly Glu Phe Arg		
215	220	
225		
aat gaa gga aac ttg gaa aag ttt gac aaa tct gct cta gag ggc ctg	720	
Asn Glu Gly Asn Leu Glu Lys Phe Asp Lys Ser Ala Leu Glu Gly Leu		
230	235	240
245		
tgc aat ttg acc att gaa gaa ttc cga tta gca tac tta gac tac tac	768	
Cys Asn Leu Thr Ile Glu Glu Phe Arg Leu Ala Tyr Leu Asp Tyr Tyr		
250	255	
260		
ctc gat gat att att gac tta ttt aat tgt ttg aca aat gtt tct tca	816	
Leu Asp Asp Ile Ile Asp Leu Phe Asn Cys Leu Thr Asn Val Ser Ser		
265	270	
275		
ttt tcc ctg gtg agt gtg act att gaa agg gta aaa gac ttt tct tat	864	
Phe Ser Leu Val Ser Val Thr Ile Glu Arg Val Lys Asp Phe Ser Tyr		
280	285	
290		
aat ttc gga tgg caa cat tta gaa tta gtt aac tgt aaa ttt gga cag	912	
Asn Phe Gly Trp Gln His Leu Glu Leu Val Asn Cys Lys Phe Gly Gln		
295	300	
305		
ttt ccc aca ttg aaa ctc aaa tct ctc aaa agg ctt act ttc act tcc	960	
Phe Pro Thr Leu Lys Leu Lys Ser Leu Lys Arg Leu Thr Phe Thr Ser		
310	315	320
325		
aac aaa ggt ggg aat gct ttt tca gaa gtt gat cta cca agc ctt gag	1008	
Asn Lys Gly Gly Asn Ala Phe Ser Glu Val Asp Leu Pro Ser Leu Glu		
330	335	
340		
ttt cta gat ctc agt aga aat ggc ttg agt ttc aaa ggt tgc tgt tct	1056	
Phe Leu Asp Leu Ser Arg Asn Gly Leu Ser Phe Lys Gly Cys Cys Ser		
345	350	
355		
caa agt gat ttt ggg aca acc agc cta aag tat tta gat ctg agc ttc	1104	
Gln Ser Asp Phe Gly Thr Thr Ser Leu Lys Tyr Leu Asp Leu Ser Phe		
360	365	
370		
aat ggt gtt att acc atg agt tca aac ttc ttg ggc tta gaa caa cta	1152	
Asn Gly Val Ile Thr Met Ser Ser Asn Phe Leu Gly Leu Glu Gln Leu		
375	380	
385		
gaa cat ctg gat ttc cag cat tcc aat ttg aaa caa atg agt gag ttt	1200	

Sequence Data File for PCT.app.txt

Glu 385	His	Leu	Asp	Phe	Gln 390	His	Ser	Asn	Leu	Lys 395	Gln	Met	Ser	Glu	Phe 400	
tca Ser	gta Val	ttc Phe	cta Leu	tca Ser 405	ctc Leu	aga Arg	aac Asn	ctc Leu	att Ile 410	tac Tyr	ctt Leu	gac Asp	att Ile	tct Ser 415	cat His	1248
act Thr	cac His	acc Thr	aga Arg 420	gtt Val	gct Ala	ttc Phe	aat Asn	ggc Gly 425	atc Ile	ttc Phe	aat Asn	ggc Gly 430	ttg Leu	tcc Ser	agt Ser	1296
ctc Leu	gaa Glu	gtc Val 435	ttg Leu	aaa Lys	atg Met	gct Ala	ggc Gly 440	aat Asn	tct Ser	ttc Phe	cag Gln	gaa Glu 445	aac Asn	ttc Phe	ctt Leu	1344
cca Pro	gat Asp 450	atc Ile	ttc Phe	aca Thr	gag Glu 455	ctg Leu	aga Arg	aac Asn	ttg Leu	acc Thr	ttc Phe 460	ctg Leu	gac Asp	ctc Leu	tct Ser	1392
cag Gln 465	tgt Cys	caa Gln	ctg Leu	gag Glu	cag Gln 470	ttg Leu	tct Ser	cca Pro	aca Thr	gca Ala 475	ttt Phe	aac Asn	tca Ser	ctc Leu	tcc Ser 480	1440
agt Ser	ctt Leu	cag Gln	gta Val	cta Leu 485	aat Asn	atg Met	agc Ser	cac His	aac Asn 490	aac Asn	ttc Phe	ttt Phe	tca Ser	ttg Leu 495	gat Asp	1488
acg Thr	ttt Phe	cct Pro	tat Tyr 500	aag Lys	tgt Cys	ctg Leu	aac Asn	tcc Ser 505	ctc Leu	cag Gln	gtt Val	ctt Leu	gat Asp 510	tac Tyr	agt Ser	1536
ctc Leu	aat Asn	cac His 515	ata Ile	atg Met	act Thr	tcc Ser	aaa Lys 520	aaa Lys	cag Gln	gaa Glu	cta Leu	cag Gln 525	cat His	ttt Phe	cca Pro	1584
agt Ser	agt Ser 530	cta Leu	gct Ala	ttc Phe	tta Leu	aat Asn 535	ctt Leu	act Thr	cag Gln	aat Asn	gac Asp 540	ttt Phe	gct Ala	tgt Cys	act Thr	1632
tgt Cys 545	gaa Glu	cac His	cag Gln	agt Ser	ttc Phe 550	ctg Leu	caa Gln	tgg Trp	atc Ile	aag Lys 555	gac Asp	cag Gln	agg Arg	cag Gln	ctc Leu 560	1680
ttg Leu	gtg Val	gaa Glu	gtt Val	gaa Glu 565	cga Arg	atg Met	gaa Glu	tgt Cys	gca Ala 570	aca Thr	cct Pro	tca Ser	gat Asp	aag Lys 575	cag Gln	1728
ggc Gly	atg Met	cct Pro	gtg Val 580	ctg Leu	agt Ser	ttg Leu	aat Asn	atc Ile 585	acc Thr	tgt Cys	cag Gln	atg Met	aat Asn 590	aag Lys	acc Thr	1776
atc Ile	att Ile	ggc Gly 595	gtg Val	tcg Ser	gtc Val	ctc Leu	agt Ser 600	gtg Val	ctt Leu	gta Val	gta Val	tct Ser 605	gtt Val	gta Val	gca Ala	1824
gtt Val 610	ctg Leu	gtc Val	tat Tyr	aag Lys	ttc Phe	tat Tyr 615	ttt Phe	cac His	ctg Leu	atg Met	ctt Leu 620	ctt Leu	gct Ala	ggc Gly	tgc Cys	1872
ata Ile 625	aag Lys	tat Tyr	ggc Gly	aga Arg	ggc Gly 630	gaa Glu	aac Asn	atc Ile	tat Tyr	gat Asp 635	gcc Ala	ttt Phe	gtt Val	atc Ile	tac Tyr 640	1920

Sequence Data File for PCT.app.txt

tca	agc	cag	gat	gag	gac	tgg	gta	agg	aat	gag	cta	gta	aag	aat	tta	1968
Ser	Ser	Gln	Asp	Glu	Asp	Trp	Val	Arg	Asn	Glu	Leu	Val	Lys	Asn	Leu	
				645					650					655		
gaa	gaa	ggg	gtg	cct	cca	ttt	cag	ctc	tgc	ctt	cac	tac	aga	gac	ttt	2016
Glu	Glu	Gly	Val	Pro	Pro	Phe	Gln	Leu	Cys	Leu	His	Tyr	Arg	Asp	Phe	
			660					665					670			
att	ccc	ggg	gtg	gcc	att	gct	gcc	aac	atc	atc	cat	gaa	ggg	ttc	cat	2064
Ile	Pro	Gly	Val	Ala	Ile	Ala	Ala	Asn	Ile	Ile	His	Glu	Gly	Phe	His	
		675					680					685				
aaa	agc	cga	aag	gtg	att	gtt	gtg	gtg	tcc	cag	cac	ttc	atc	cag	agc	2112
Lys	Ser	Arg	Lys	Val	Ile	Val	Val	Val	Ser	Gln	His	Phe	Ile	Gln	Ser	
	690					695					700					
cgc	tgg	tgt	atc	ttt	gaa	tat	gag	att	gct	cag	acc	tgg	cag	ttt	ctg	2160
Arg	Trp	Cys	Ile	Phe	Glu	Tyr	Glu	Ile	Ala	Gln	Thr	Trp	Gln	Phe	Leu	
705					710					715					720	
agc	agt	cgt	gct	ggg	atc	atc	ttc	att	gtc	ctg	cag	aag	gtg	gag	aag	2208
Ser	Ser	Arg	Ala	Gly	Ile	Ile	Phe	Ile	Val	Leu	Gln	Lys	Val	Glu	Lys	
				725					730					735		
acc	ctg	ctc	agg	cag	cag	gtg	gag	ctg	tac	cgc	ctt	ctc	agc	agg	aac	2256
Thr	Leu	Leu	Arg	Gln	Gln	Val	Glu	Leu	Tyr	Arg	Leu	Leu	Ser	Arg	Asn	
			740					745					750			
act	tac	ctg	gag	tgg	gag	gac	agt	gtc	ctg	ggg	cgg	cac	atc	ttc	tgg	2304
Thr	Tyr	Leu	Glu	Trp	Glu	Asp	Ser	Val	Leu	Gly	Arg	His	Ile	Phe	Trp	
		755					760					765				
aga	cga	ctc	aga	aaa	gcc	ctg	ctg	gat	ggg	aaa	tca	tgg	aat	cca	gaa	2352
Arg	Arg	Leu	Arg	Lys	Ala	Leu	Leu	Asp	Gly	Lys	Ser	Trp	Asn	Pro	Glu	
	770					775					780					
gga	aca	gtg	ggg	aca	gga	tgc	aat	tgg	cag	gaa	gca	aca	tct	atc	tga	2400
Gly	Thr	Val	Gly	Thr	Gly	Cys	Asn	Trp	Gln	Glu	Ala	Thr	Ser	Ile		
785					790					795						

<210> 7
 <211> 799
 <212> PRT
 <213> Homo sapiens

<400> 7
 Met Glu Leu Asn Phe Tyr Lys Ile Pro Asp Asn Leu Pro Phe Ser Thr
 1 5 10 15
 Lys Asn Leu Asp Leu Ser Phe Asn Pro Leu Arg His Leu Gly Ser Tyr
 20 25 30
 Ser Phe Phe Ser Phe Pro Glu Leu Gln Val Leu Asp Leu Ser Arg Cys
 35 40 45
 Glu Ile Gln Thr Ile Glu Asp Gly Ala Tyr Gln Ser Leu Ser His Leu
 50 55 60
 Ser Thr Leu Ile Leu Thr Gly Asn Pro Ile Gln Ser Leu Ala Leu Gly
 65 70 75 80
 Ala Phe Ser Gly Leu Ser Ser Leu Gln Lys Leu Val Ala Val Glu Thr
 Page 5

Sequence Data File for PCT.app.txt

85

90

95

Asn Leu Ala Ser Leu Glu Asn Phe Pro Ile Gly His Leu Lys Thr Leu
 100 105 110
 Lys Glu Leu Asn Val Ala His Asn Leu Ile Gln Ser Phe Lys Leu Pro
 115 120 125
 Glu Tyr Phe Ser Asn Leu Thr Asn Leu Glu His Leu Asp Leu Ser Ser
 130 135 140
 Asn Lys Ile Gln Ser Ile Tyr Cys Thr Asp Leu Arg Val Leu His Gln
 145 150 155 160
 Met Pro Leu Leu Asn Leu Ser Leu Asp Leu Ser Leu Asn Pro Met Asn
 165 170 175
 Phe Ile Gln Pro Gly Ala Phe Lys Glu Ile Arg Leu His Lys Leu Thr
 180 185 190
 Leu Arg Asn Asn Phe Asp Ser Leu Asn Val Met Lys Thr Cys Ile Gln
 195 200 205
 Gly Leu Ala Gly Leu Glu Val His Arg Leu Val Leu Gly Glu Phe Arg
 210 215 220
 Asn Glu Gly Asn Leu Glu Lys Phe Asp Lys Ser Ala Leu Glu Gly Leu
 225 230 235 240
 Cys Asn Leu Thr Ile Glu Glu Phe Arg Leu Ala Tyr Leu Asp Tyr Tyr
 245 250 255
 Leu Asp Asp Ile Ile Asp Leu Phe Asn Cys Leu Thr Asn Val Ser Ser
 260 265 270
 Phe Ser Leu Val Ser Val Thr Ile Glu Arg Val Lys Asp Phe Ser Tyr
 275 280 285
 Asn Phe Gly Trp Gln His Leu Glu Leu Val Asn Cys Lys Phe Gly Gln
 290 295 300
 Phe Pro Thr Leu Lys Leu Lys Ser Leu Lys Arg Leu Thr Phe Thr Ser
 305 310 315 320
 Asn Lys Gly Gly Asn Ala Phe Ser Glu Val Asp Leu Pro Ser Leu Glu
 325 330 335
 Phe Leu Asp Leu Ser Arg Asn Gly Leu Ser Phe Lys Gly Cys Cys Ser
 340 345 350
 Gln Ser Asp Phe Gly Thr Thr Ser Leu Lys Tyr Leu Asp Leu Ser Phe
 355 360 365
 Asn Gly Val Ile Thr Met Ser Ser Asn Phe Leu Gly Leu Glu Gln Leu
 370 375 380
 Glu His Leu Asp Phe Gln His Ser Asn Leu Lys Gln Met Ser Glu Phe
 385 390 395 400
 Ser Val Phe Leu Ser Leu Arg Asn Leu Ile Tyr Leu Asp Ile Ser His
 405 410 415
 Thr His Thr Arg Val Ala Phe Asn Gly Ile Phe Asn Gly Leu Ser Ser

Sequence Data File for PCT.app.txt

420
 Leu Glu Val Leu Lys Met Ala Gly Asn Ser Phe Gln Glu Asn Phe Leu
 435 440 445
 Pro Asp Ile Phe Thr Glu Leu Arg Asn Leu Thr Phe Leu Asp Leu Ser
 450 455 460
 Gln Cys Gln Leu Glu Gln Leu Ser Pro Thr Ala Phe Asn Ser Leu Ser
 465 470 475 480
 Ser Leu Gln Val Leu Asn Met Ser His Asn Asn Phe Phe Ser Leu Asp
 485 490 495
 Thr Phe Pro Tyr Lys Cys Leu Asn Ser Leu Gln Val Leu Asp Tyr Ser
 500 505 510
 Leu Asn His Ile Met Thr Ser Lys Lys Gln Glu Leu Gln His Phe Pro
 515 520 525
 Ser Ser Leu Ala Phe Leu Asn Leu Thr Gln Asn Asp Phe Ala Cys Thr
 530 535 540
 Cys Glu His Gln Ser Phe Leu Gln Trp Ile Lys Asp Gln Arg Gln Leu
 545 550 555 560
 Leu Val Glu Val Glu Arg Met Glu Cys Ala Thr Pro Ser Asp Lys Gln
 565 570 575
 Gly Met Pro Val Leu Ser Leu Asn Ile Thr Cys Gln Met Asn Lys Thr
 580 585 590
 Ile Ile Gly Val Ser Val Leu Ser Val Leu Val Val Ser Val Val Ala
 595 600 605
 Val Leu Val Tyr Lys Phe Tyr Phe His Leu Met Leu Leu Ala Gly Cys
 610 615 620
 Ile Lys Tyr Gly Arg Gly Glu Asn Ile Tyr Asp Ala Phe Val Ile Tyr
 625 630 635 640
 Ser Ser Gln Asp Glu Asp Trp Val Arg Asn Glu Leu Val Lys Asn Leu
 645 650 655
 Glu Glu Gly Val Pro Pro Phe Gln Leu Cys Leu His Tyr Arg Asp Phe
 660 665 670
 Ile Pro Gly Val Ala Ile Ala Ala Asn Ile Ile His Glu Gly Phe His
 675 680 685
 Lys Ser Arg Lys Val Ile Val Val Val Ser Gln His Phe Ile Gln Ser
 690 695 700
 Arg Trp Cys Ile Phe Glu Tyr Glu Ile Ala Gln Thr Trp Gln Phe Leu
 705 710 715 720
 Ser Ser Arg Ala Gly Ile Ile Phe Ile Val Leu Gln Lys Val Glu Lys
 725 730 735
 Thr Leu Leu Arg Gln Gln Val Glu Leu Tyr Arg Leu Leu Ser Arg Asn
 740 745 750
 Thr Tyr Leu Glu Trp Glu Asp Ser Val Leu Gly Arg His Ile Phe Trp

Sequence Data File for PCT.app.txt
755 760 765
Arg Arg Leu Arg Lys Ala Leu Leu Asp Gly Lys Ser Trp Asn Pro Glu
770 775 780
Gly Thr Val Gly Thr Gly Cys Asn Trp Gln Glu Ala Thr Ser Ile
785 790 795

INTERNATIONAL SEARCH REPORT

Inte bnal Application No
PCT/US 02/34120

A. CLASSIFICATION OF SUBJECT MATTER

IPC 7 A61K45/00 A61L31/16 A61P9/10 A61K31/7105 A61K38/17
A61K39/395

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 A61K A61L A61P

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

WPI Data, EPO-Internal, PAJ, MEDLINE, BIOSIS, CHEM ABS Data, SCISEARCH

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 00 72023 A (SINGH MAHAVIR ;FUST GYORGY (HU); ROMICS LASZLO (HU); PROHASZKA ZOL) 30 November 2000 (2000-11-30)	29-60
Y	page 8, line 14 - line 24	1-28
X	WO 01 43691 A (KOLB HUBERT ;ELIAS DANA (IL); PEPTOR LTD (IL)) 21 June 2001 (2001-06-21)	29-60
Y	page 6, line 13 - line 16 page 12, line 9 - line 15; claims 19,20	1-28
X	WO 00 20019 A (HARATS DROR ;GEORGE JACOB (IL); SHOENFELD YEHUDA (IL)) 13 April 2000 (2000-04-13)	29-60
Y	page 10, line 20 - line 26	1-28
	--- -/--	



Further documents are listed in the continuation of box C.



Patent family members are listed in annex.

* Special categories of cited documents :

A document defining the general state of the art which is not considered to be of particular relevance

E earlier document but published on or after the international filing date

L document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

O document referring to an oral disclosure, use, exhibition or other means

P document published prior to the international filing date but later than the priority date claimed

T later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

X document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

Y document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

G document member of the same patent family

Date of the actual completion of the international search

19 February 2003

Date of mailing of the international search report

28/02/2003

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2
NL - 2280 HV Rijswijk
Tel. (+31-70) 340-2040, Tx. 31 651 epo nl,
Fax: (+31-70) 340-3016

Authorized officer

Pilling, S

INTERNATIONAL SEARCH REPORT

International Application No
PCT/US 02/34120

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	US 5 733 327 A (IGAKI KEIJI ET AL) 31 March 1998 (1998-03-31) column 2, line 54 -column 3, line 18; examples 1-7	1-28
Y	US 5 591 227 A (SCHWARTZ ROBERT S ET AL) 7 January 1997 (1997-01-07) column 2, line 36 - line 56	1-28
Y	US 6 171 609 B1 (KUNZ LAWRENCE L) 9 January 2001 (2001-01-09) column 4, line 8 - line 43; claim 1	1-28
A	XIAOOU XU HELEN ET AL: "Toll-Like Receptor-4 is expressed in human coronary atherosclerotic plaques and upregulated by oxidized low density lipoprotein in macrophages" JOURNAL OF THE AMERICAN COLLEGE OF CARDIOLOGY, XX, XX, vol. 35, no. 2 suppl A, 1 February 2000 (2000-02-01), page 260A XP002210581 ISSN: 0735-1097 abstract	1-60

INTERNATIONAL SEARCH REPORT
Information on patent family members

International Application No
PCT/US 02/34120

Patent document cited in search report		Publication date	Patent family member(s)	Publication date
WO 0072023	A	30-11-2000	AU 4602500 A	12-12-2000
			BR 0010741 A	19-02-2002
			CN 1351713 T	29-05-2002
			EP 1179182 A2	13-02-2002
			HU 0201568 A2	28-08-2002
			WO 0072023 A2	30-11-2000
			NO 20015653 A	17-01-2002
WO 0143691	A	21-06-2001	AU 1881601 A	25-06-2001
			EP 1237570 A2	11-09-2002
			WO 0143691 A2	21-06-2001
WO 0020019	A	13-04-2000	AU 5996699 A	26-04-2000
			BR 9914631 A	03-07-2001
			CA 2345445 A1	13-04-2000
			CN 1332636 T	23-01-2002
			EP 1126867 A2	29-08-2001
			HU 0104410 A2	28-03-2002
			WO 0020019 A2	13-04-2000
			JP 2002526416 T	20-08-2002
			US 2002025321 A1	28-02-2002
US 5733327	A	31-03-1998	AU 699821 B2	17-12-1998
			AU 3674295 A	06-05-1996
			EP 1181904 A2	27-02-2002
			EP 0761251 A1	12-03-1997
			WO 9611720 A1	25-04-1996
US 5591227	A	07-01-1997	US 5599352 A	04-02-1997
			US 5957971 A	28-09-1999
			DE 69527900 D1	02-10-2002
			EP 0701802 A1	20-03-1996
			JP 8089585 A	09-04-1996
			US 5697967 A	16-12-1997
			US 6080190 A	27-06-2000
			US 5571166 A	05-11-1996
			US 5591224 A	07-01-1997
			US 5510077 A	23-04-1996
			US 5554182 A	10-09-1996
			US 5800507 A	01-09-1998
			US 5628785 A	13-05-1997
			US 5849034 A	15-12-1998
			DE 69326631 D1	11-11-1999
			DE 69326631 T2	08-06-2000
			EP 0566245 A1	20-10-1993
			JP 6007455 A	18-01-1994
US 6171609	B1	09-01-2001	US 6515009 B1	04-02-2003
			AU 4985196 A	04-09-1996
			CA 2212537 A1	22-08-1996
			EP 0809515 A1	03-12-1997
			JP 11500635 T	19-01-1999
			US 2002086896 A1	04-07-2002
			WO 9625176 A1	22-08-1996
			US 6358989 B1	19-03-2002
			US 5981568 A	09-11-1999
			US 6306421 B1	23-10-2001
			US 2002025979 A1	28-02-2002

INTERNATIONAL SEARCH REPORT
Information on patent family members

International Application No
PCT/US 02/34120

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
US 6171609	B1	US 2002040064 A1	04-04-2002